-continued

<213> ORGANISM: Diabrotica virgifera

<400> SEQUENCE: 131

ctagtggtga tatatatggt ctagttcatg ag 32

<210> SEQ ID NO 132
<211> LENGTH: 32
<212> TYPE: DNA
<213> ORGANISM: Diabrotica balteata

<400> SEQUENCE: 132

aattctcatg aactagacca tatatatcac ca 32

<210> SEQ ID NO 133
<211> LENGTH: 32
<212> TYPE: DNA
<213> ORGANISM: Diabrotica virgifera zeae

<400> SEQUENCE: 133

ctagttatat atggtctagt tcatgaaaac ag 32

<210> SEQ ID NO 134
<211> LENGTH: 32
<212> TYPE: DNA
<213> ORGANISM: Diabrotica balteata

<400> SEQUENCE: 134

aattctgttt tcatgaacta gaccatatat aa 32

<210> SEQ ID NO 135
<211> LENGTH: 32
<212> TYPE: DNA
<213> ORGANISM: Hippodamia convergens

<400> SEQUENCE: 135

ctagtatggt ctagttcatg aaaacaccct tg 32

1-95. (canceled)

96. A method for managing crop pest resistance to an agent for controlling the crop pest, comprising contacting the crop pest with a nucleic acid segment produced by a method comprising:

(a) obtaining a starting nucleic acid molecule substantially complementary to a target gene;

(b) preparing a plurality of nucleic acid segments from the starting nucleic acid molecule;

(c) assaying the nucleic acid segments for the ability to suppress expression of the target gene when expressed as a dsRNA in a cell comprising the target gene; and

(d) identifying at least a first nucleic acid segment from the plurality of nucleic acid segments that provides a desired level of suppression of the target gene when expressed as a dsRNA,

and one additional agent selected from the group consisting of a patatin, a *Bacillus thuringiensis* insecticidal protein, a *Xenorhabdus* insecticidal protein, a *Photorhabdus* insecticidal protein, a *Bacillus* laterosporus insecticidal protein, a *Bacillus sphaericus* insecticidal protein, a biocontrol agent, and an insecticide.

97. The method of claim **96**, wherein the insecticide is selected from the group consisting of a carbaryl insecticide, fenvalerate, esfenvalerate, malathion, a carbofuran insecticide, chloropyrifos, fonophos, phorate, terbufos, permethrin, a neonicotinoid, and tefluthrin.

98. The method of claim **96**, wherein the additional agent is provided as a seed treatment.

99. The method of claim **96** wherein the additional agent is a *Bacillus thuringiensis* insecticidal protein.

100. The method of claim **99**, wherein the *Bacillus thuringiensis* insecticidal protein is selected from the group consisting of a Cry1, a Cry3, a TIC851, a CryET70, a Cry2, ET29, ET37, a binary insecticidal protein CryET33 and CryET34, a binary insecticidal protein CryET80 and CryET76, a binary insecticidal protein TIC100 and TIC101, a binary insecticidal protein ET29 and TIC810, a binary insecticidal protein ET37 and TIC812, and a binary insecticidal protein PS 149B 1.

101. The method of claim **96**, wherein the desired level of suppression is complete suppression of the target gene.

102. The method of claim **96**, wherein the desired level of suppression is incomplete suppression of the target gene.

103. The method of claim **96**, wherein assaying the nucleic acid segments for the ability to suppress the target gene comprises expressing the segments as a dsRNA in a cell comprising the target gene and determining the level of suppression of the target gene.